



FIGURE 1

*** ATF *** SP1 *** GC
 182510 aAGGTCCACGTCAGGCGGGGCCCCGCCACCGCGGACCAAGCGGCTACGCCTTCCAAGT (SEQ ID NO:3)
 ss3217444 caccgcgggaccaagcggtacgccttccargt (SEQ ID NO:4)

*** VMAF *** TAl1alphaE47 ***
 182570 TTTCATTCCGGGCGCGCCAGCGCTGAGGAGAGTCAGACGCAGATGGTTCCGAAGGTCC (SEQ ID NO:5)
 tttcattcggggcgcgcccagcgctgag (SEQ ID NO:6)

*** AHRARNT ***
 182630 TGCAGGCTCTCGGGGCACCCTACAGCCGACACGCCAACAAAAGCTTCGGAGCCGCCAAC (SEQ ID NO:7)

*** OLF1 ***
 182690 CCAAACCGAAACACCGGAGCCACAAGCTGAGCTCGCGTAGGGCACGCTCCCCGGGAGT (SEQ ID NO:8)

*** SP1 *** NRF2 *** AP4 ***
 182750 GCTTCGGGACAGGGGCGGGGTTCGCGGAAGTGGCCGTCCGCCGTCCGCTGCTCGTTGT (SEQ ID NO:9)

*** TAXCREB *** AP2 ***
 182810 TTAAAGCGGCTGACGGGAAGGAGAAGCCAGAGCTCCAGCGGCGCCGCGGGGCGGCAGTCAA (SEQ ID NO:10)

*** PAX5 ***
 182870 GACCAGAGCCGGAGCCGTCCTACTCACCTCTGGATTAGCCTGAAGCGGAGACTACCGGCTGC (SEQ ID NO:11)

*** E47 ***
 182930 GGAGCGGCGGGGCGAGACACTTGCTCGCCTTTTGACCCCATCATGTGCGGGGCTCCATC (SEQ ID NO:12)

*** PAX5 *** SP1 ***
 182990 GAGATTCCCCTCCGGGACACTGACGAGGTAAGTGTCTGTATGGAGGCGGGTGGGGGCCA (SEQ ID NO:13)
 ss3217445 tggaggcgggtgggggcca (SEQ ID NO:14)

*** PAX5 *** PAX5 ***
 183050 TGAACCTGTACGATCGCCCCACCGACTTCGCTCGACTTCGTTTCTGAAGCGAGAGCATC (SEQ ID NO:15)
 tgaacttgtaRatcgccccaccgacttcgctcgacttcgt (SEQ ID NO:16)

**
 183110 CTccttcctaagagccctgatcgaaatacctggccaagggtctgatcatcatcttcttca (SEQ ID NO:17)

FIGURE 2